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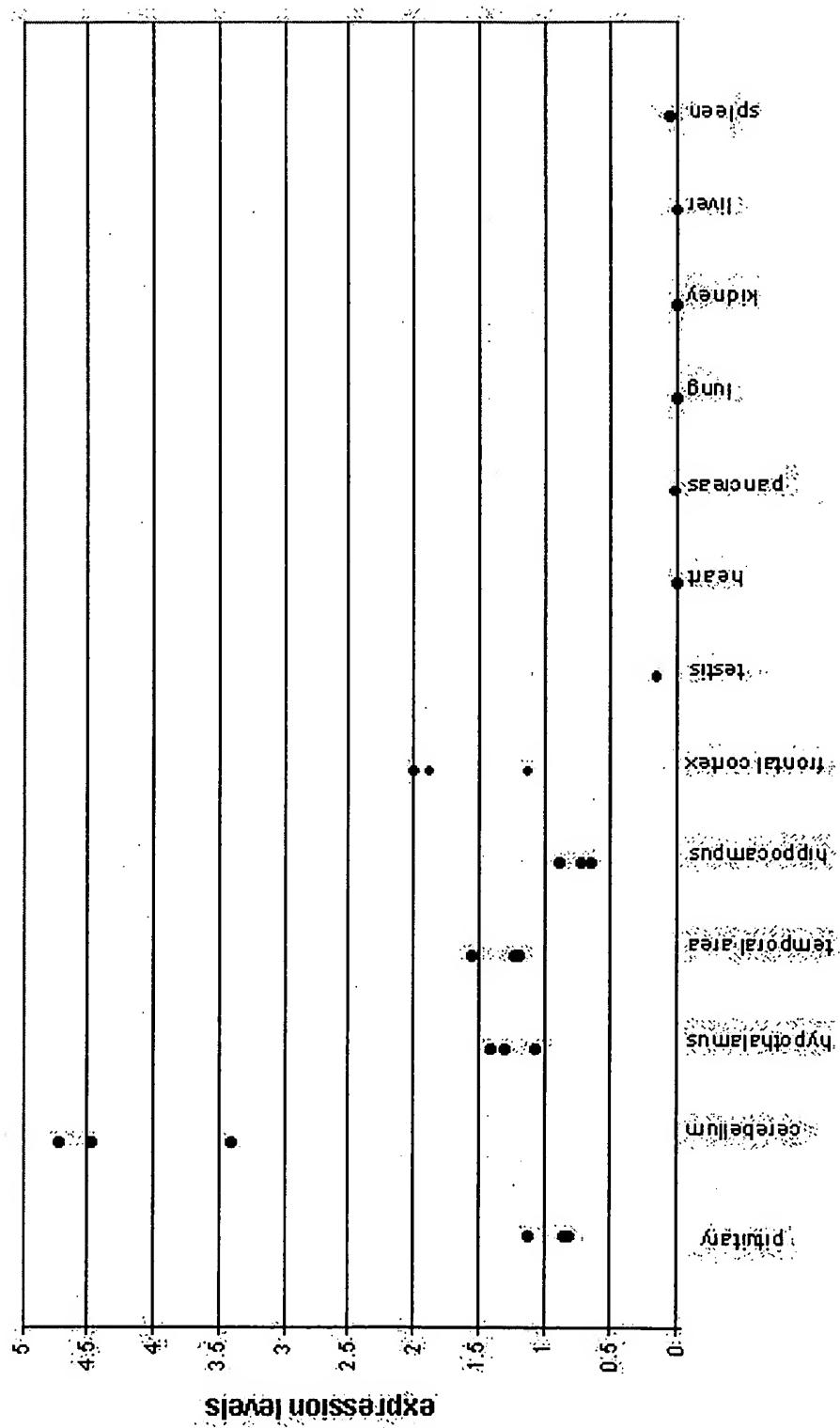
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Table 1

Gene Name	CDS	SEQ ID	NA	AA	Title	Sequence Derived from
down	1700016ED5Rik	1			RIKEN cDNA 1700016ED5 gene	AB22377
down	2300021F08Rik	2			RIKEN cDNA 2300021F08 gene	AB22354
down	2810008F04Rik	3			RIKEN cDNA 2810008F04 gene	AB45337
down	2810511G02Rik	4			RIKEN cDNA 2810511G02 gene	AB221854
down	4831430D1Rik	5			RIKEN cDNA 4831430D1 gene	AB26942
down	A1414047	6			expressed sequence AI414047	AB1125649
down	Genera:47654	7			ESTs; moderately similar to V33: mouse visinin-like protein 3 [M: musculus]	CB118
down		8			ESTs; moderately similar to V33: mouse visinin-like protein 3 [M: musculus]	AB43393
down	B.F.ABP	9			Brain Fatty acid-binding protein gene	UC48277
down	B.F.ABP	10			Brain Fatty acid-binding protein	AA481304
down	Edg2	214:1308	11		endothelial differentiation lysophosphatidic acid G protein-coupled receptor 2	U13370
down	Fam2	1:1038	12		fibroblast growth factor receptor 2	M22382
down	Gab1	24:2111	13		growth factor receptor bound protein 2-associated protein	AB08826
down	Hpcal1	26:1942	14		hippocalcin-like 1	AB085182
down	Hsd11b1	128:1006	15		hydroxysteroid 11-beta dehydrogenase 1	X832012
down	Nis12	27:1280	16		neurotensin receptor 2	U51908
down	Nis1	344:1718	17		neurotensin receptor 1	NM_01876
down	Ywhaq	21:758	18		tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, theta polypeptide	U56243
up	1:110014:22Rik	19				
up	1200008014Rik	20			RIKEN cDNA 1200008014 gene	AB22346
up	1600023:402Rik	21			RIKEN cDNA 1600023:402 gene	AB22383
up	3110003A17Rik	22			RIKEN cDNA 3110003A17 gene	AB19397
up	3110023:10Rik	23			RIKEN cDNA 3110023:10 gene	AB833425
up	AB07667	24			expressed sequence AI47657	AB19392
up	AB08050	25			expressed sequence AI480570	AB53024
up	Fkbp5	220:1600	26		FK506 binding protein 5 (61 kDa)	AB48646
up	Pdyn	1:747	27		prodynorphin	AB382264
up	Pank1	287:1406	28		preproenkephalin 1	MS5181
up	Sak	69:1384	29		serum glucocorticoid regulated kinase 1	AB006181

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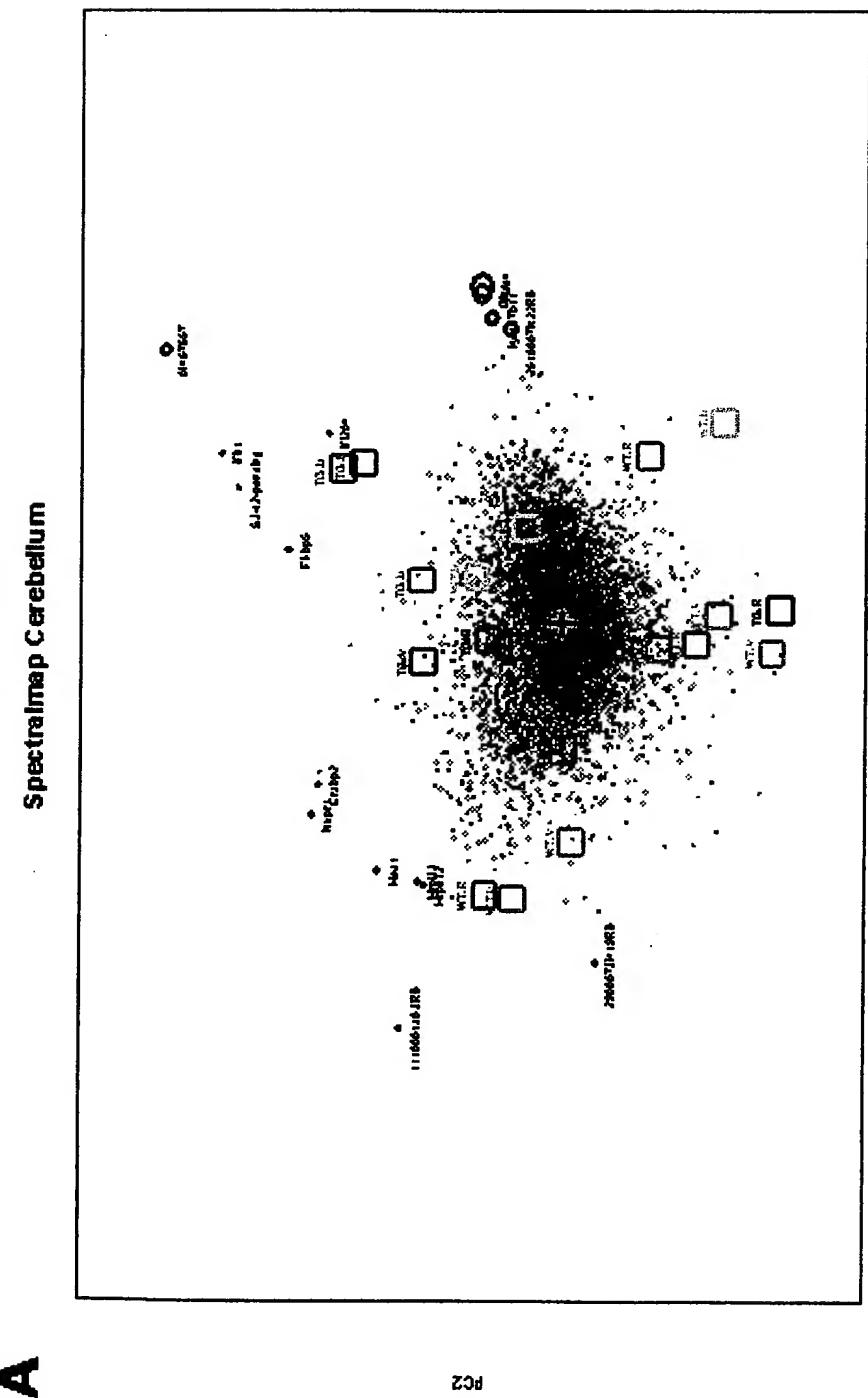
Fig. 1



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Fig. 2



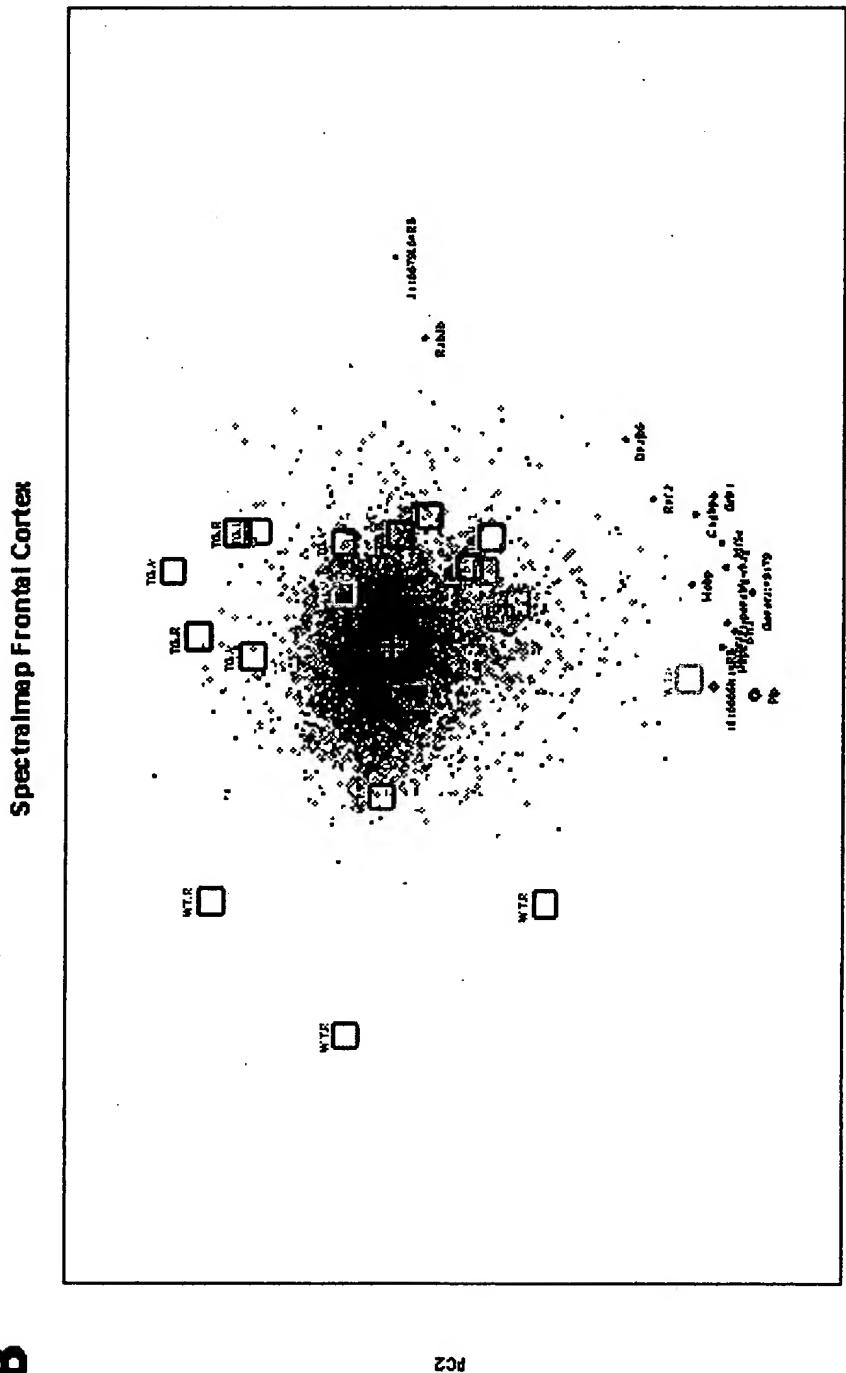
Repres. = 47% + 12%, Closure = none, Center = double, Norm. = global, Scale = WC, RUV = mean, CUV = mean

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Fig.2 continued

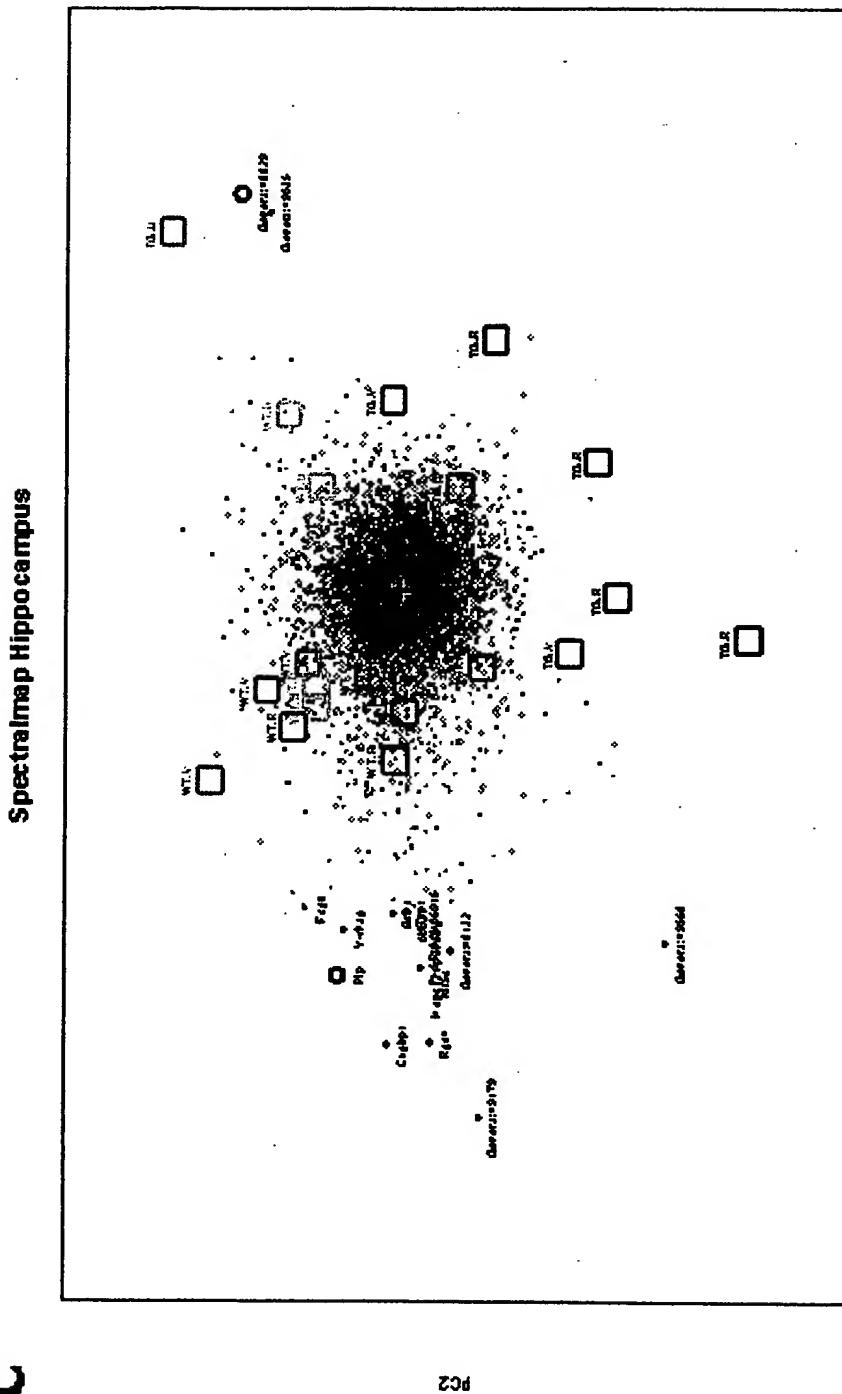


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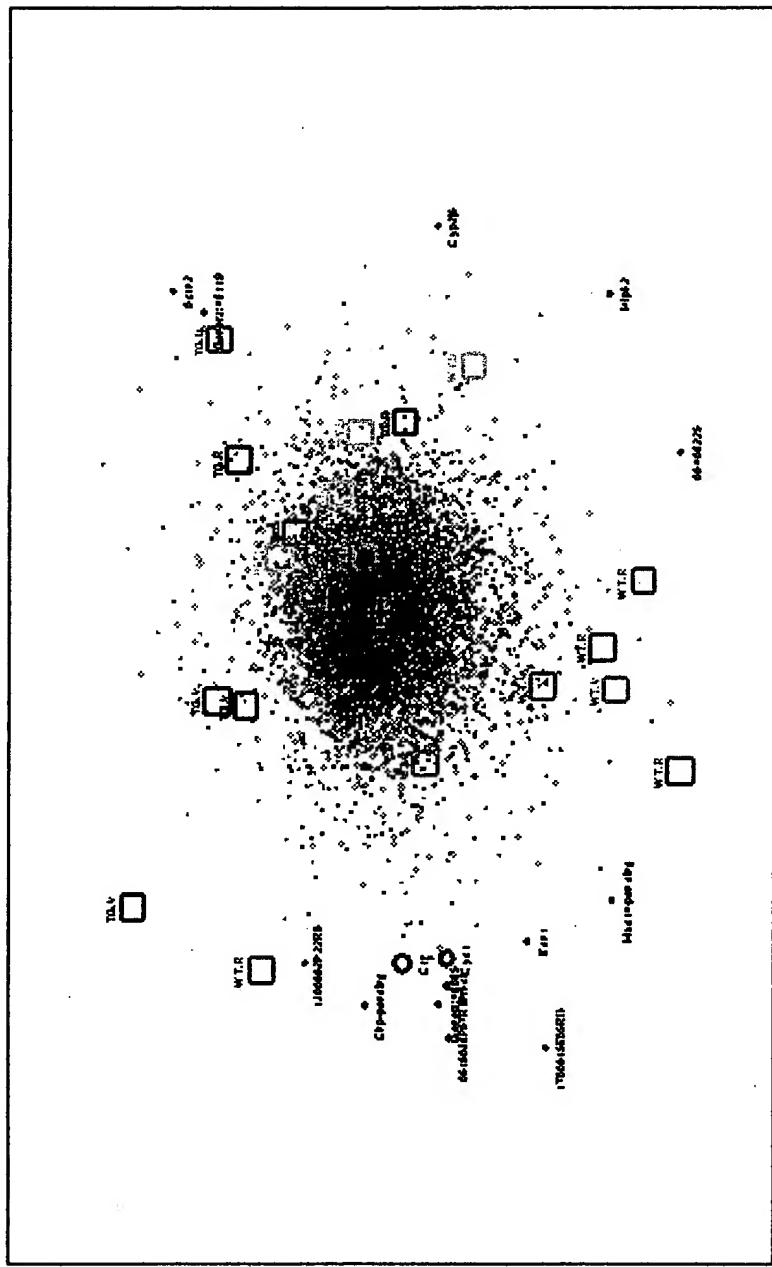
Fig.2 continued



REPLIES = 25% + 13%, CLOSURE = none, CENTER = double, NORM = global, S-CALC = uc, R00 = mean, C00 = mean

Fig.2 continued

D Spectralmap Nucleus Accumbens



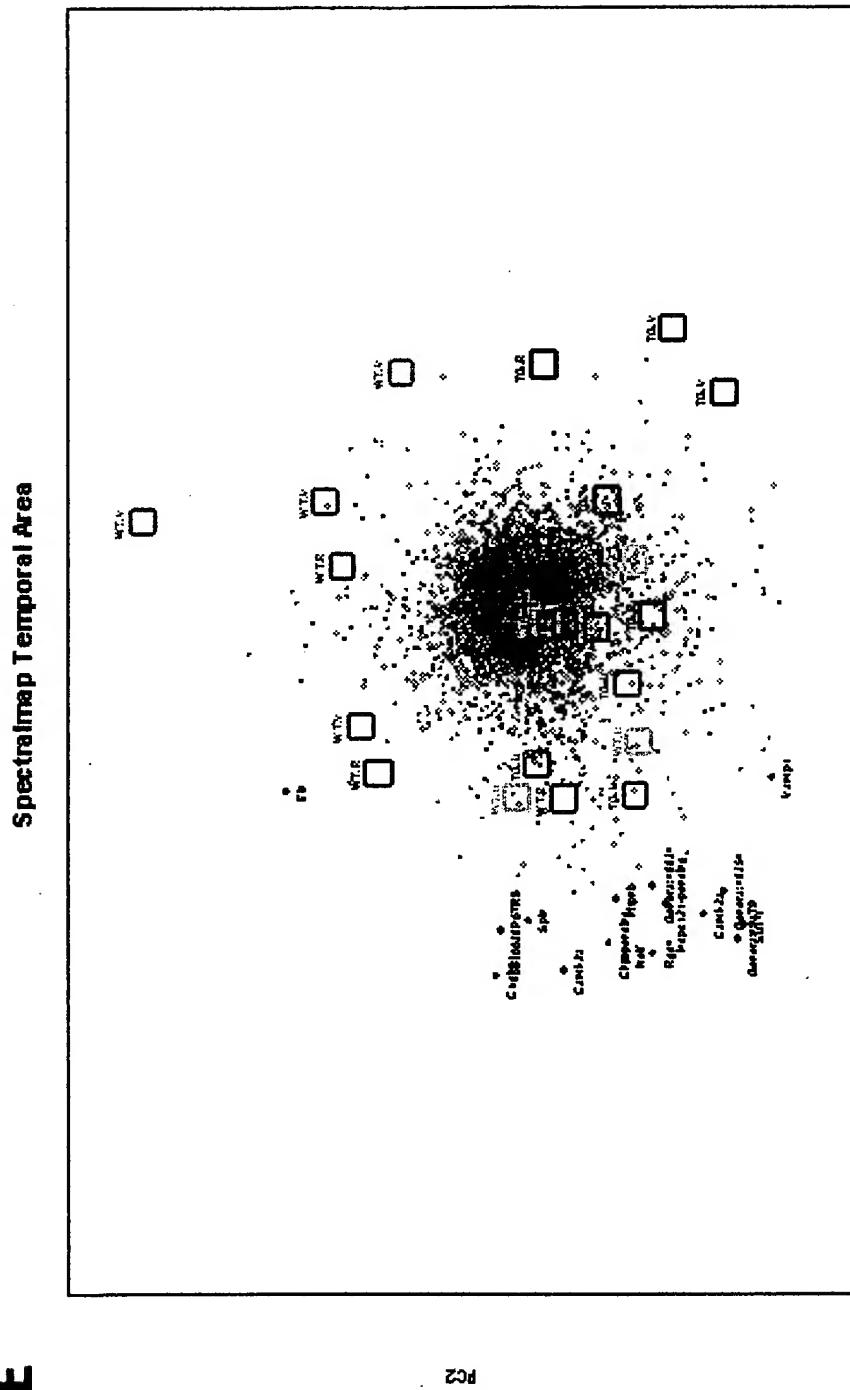
FC1
Repres. = 22% + 12%, Closure = none, Center = double, Norm. = global, Scale = 100, RUN = mean, COV = mean

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Fig.2 continued



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Fig.2 continued

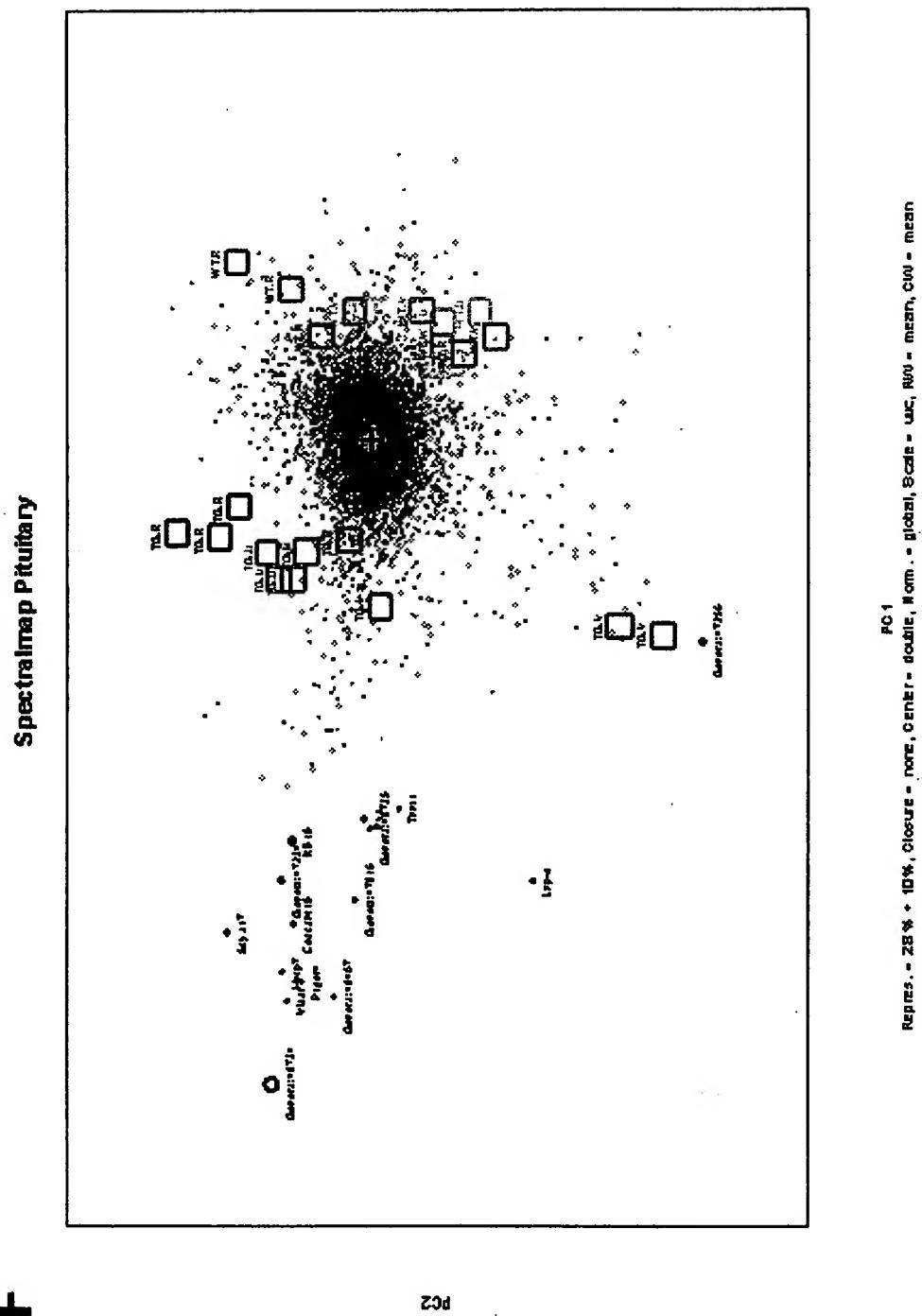
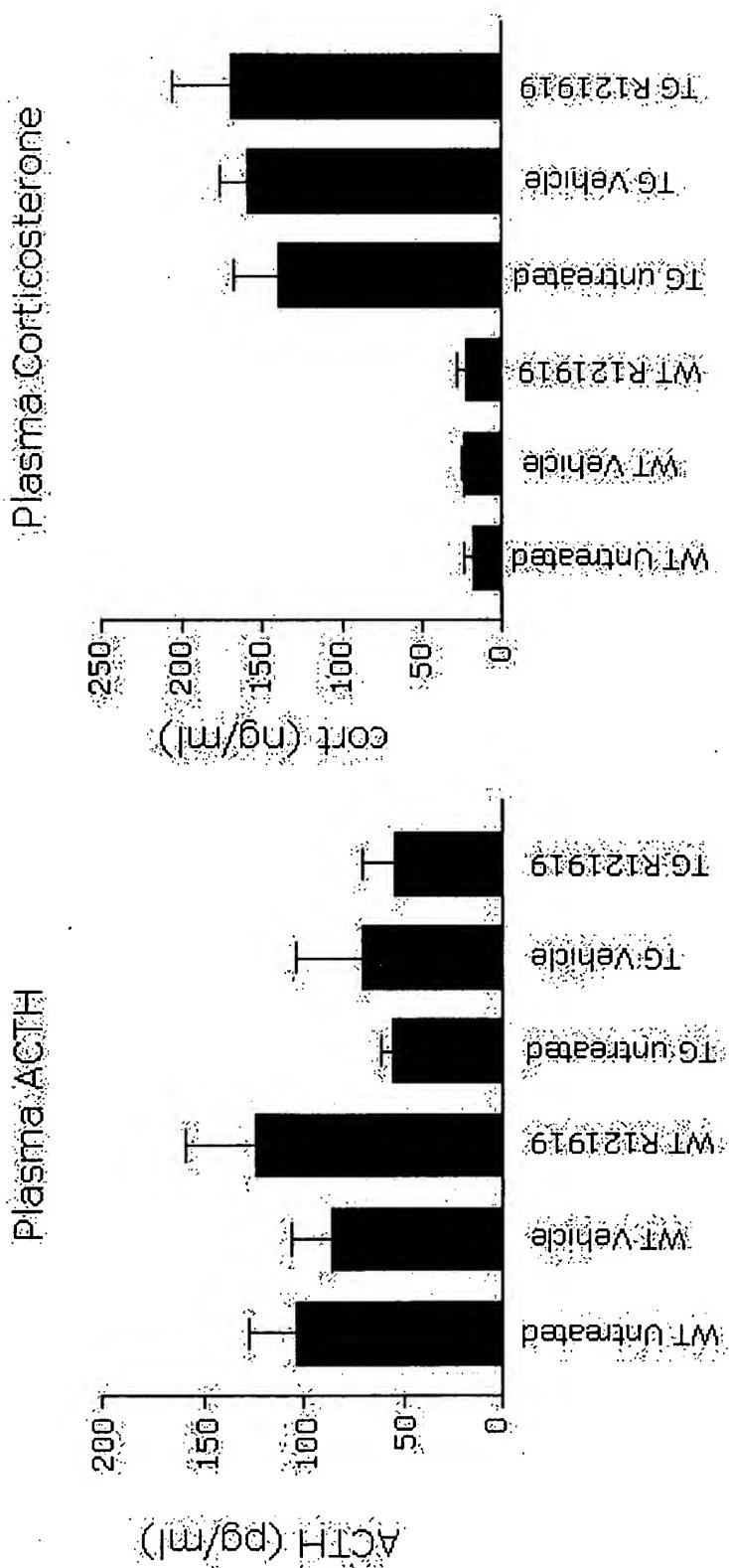


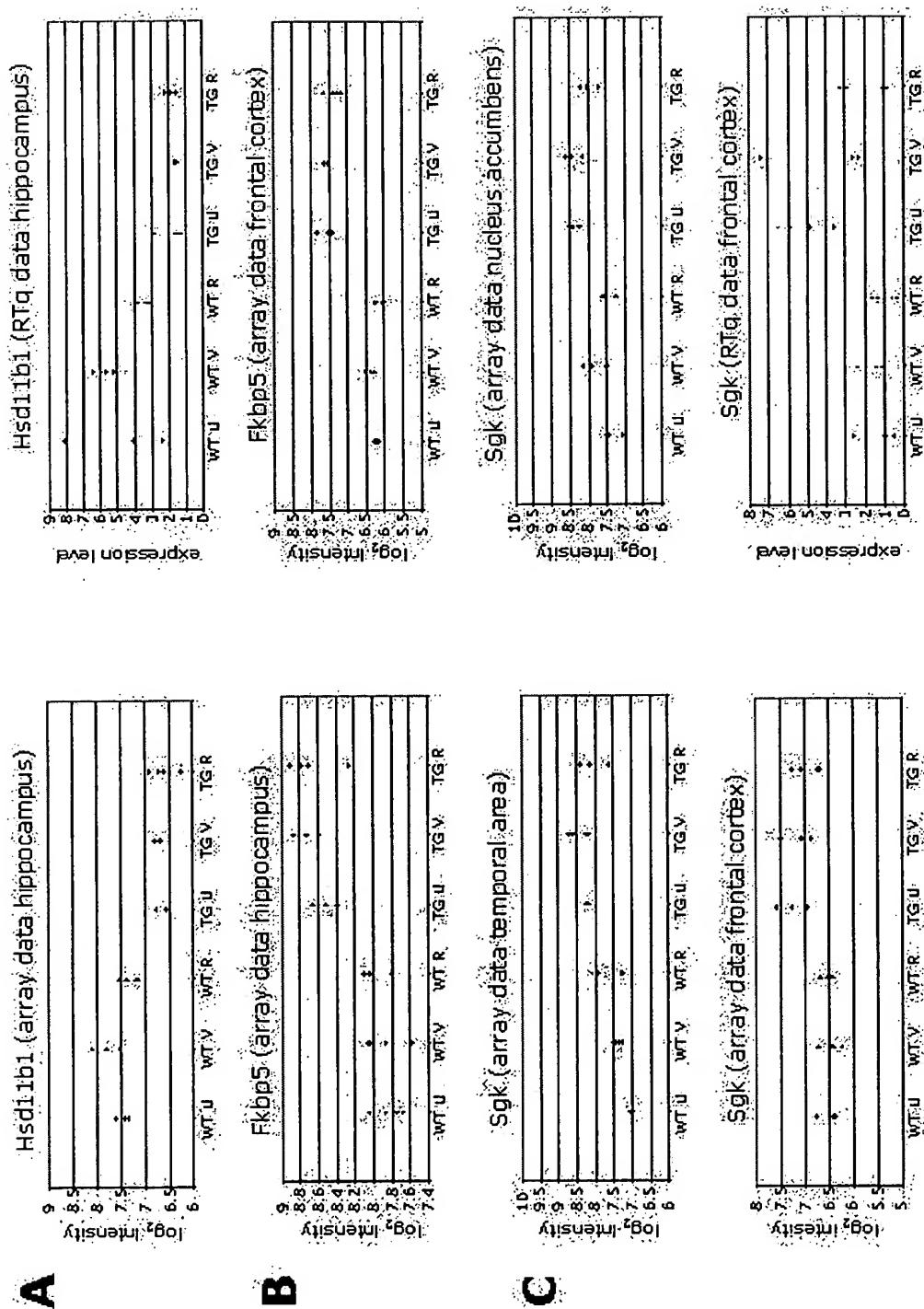
Fig. 3



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Fig. 4



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Fig.5

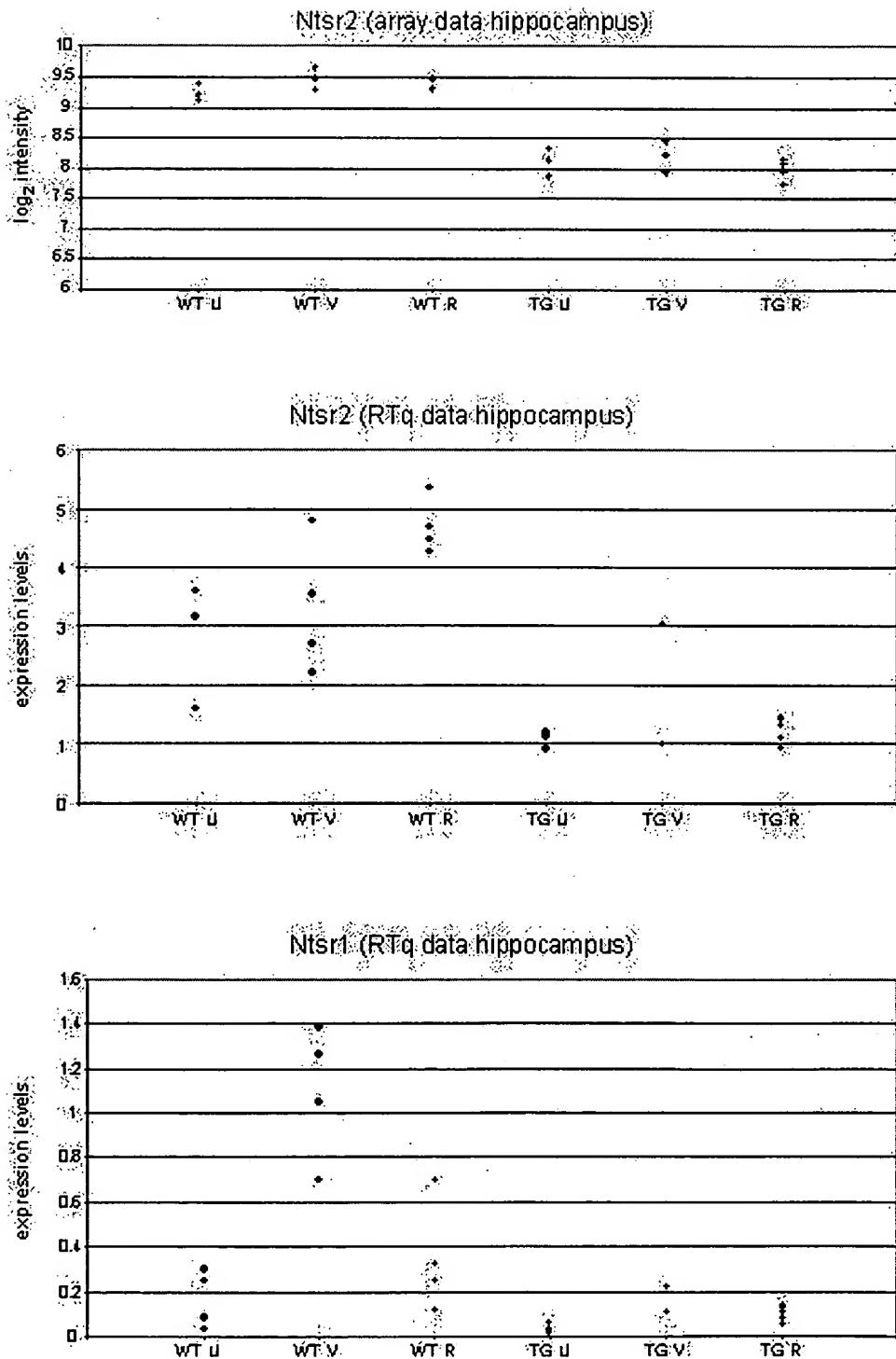


Fig.6

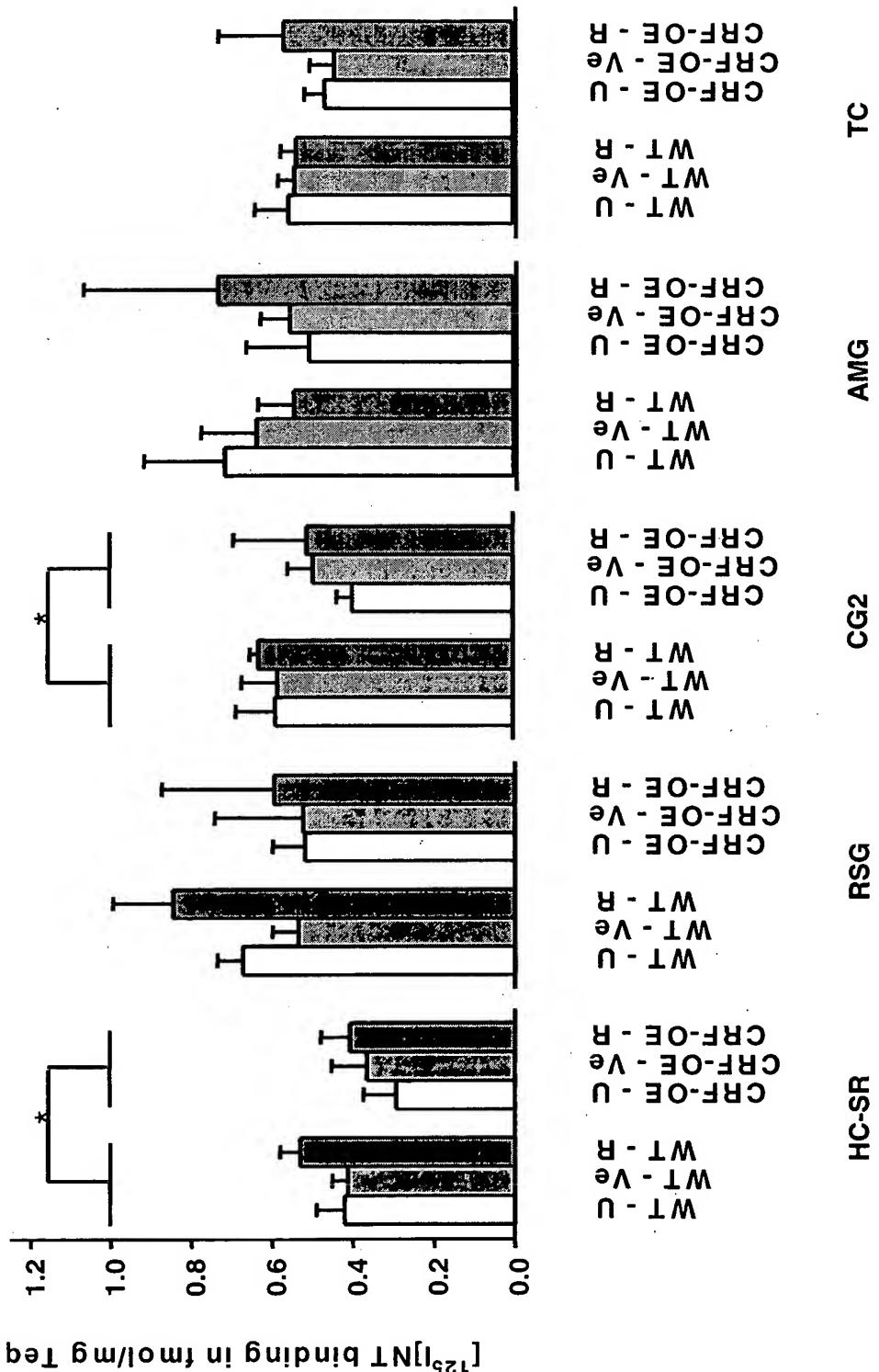
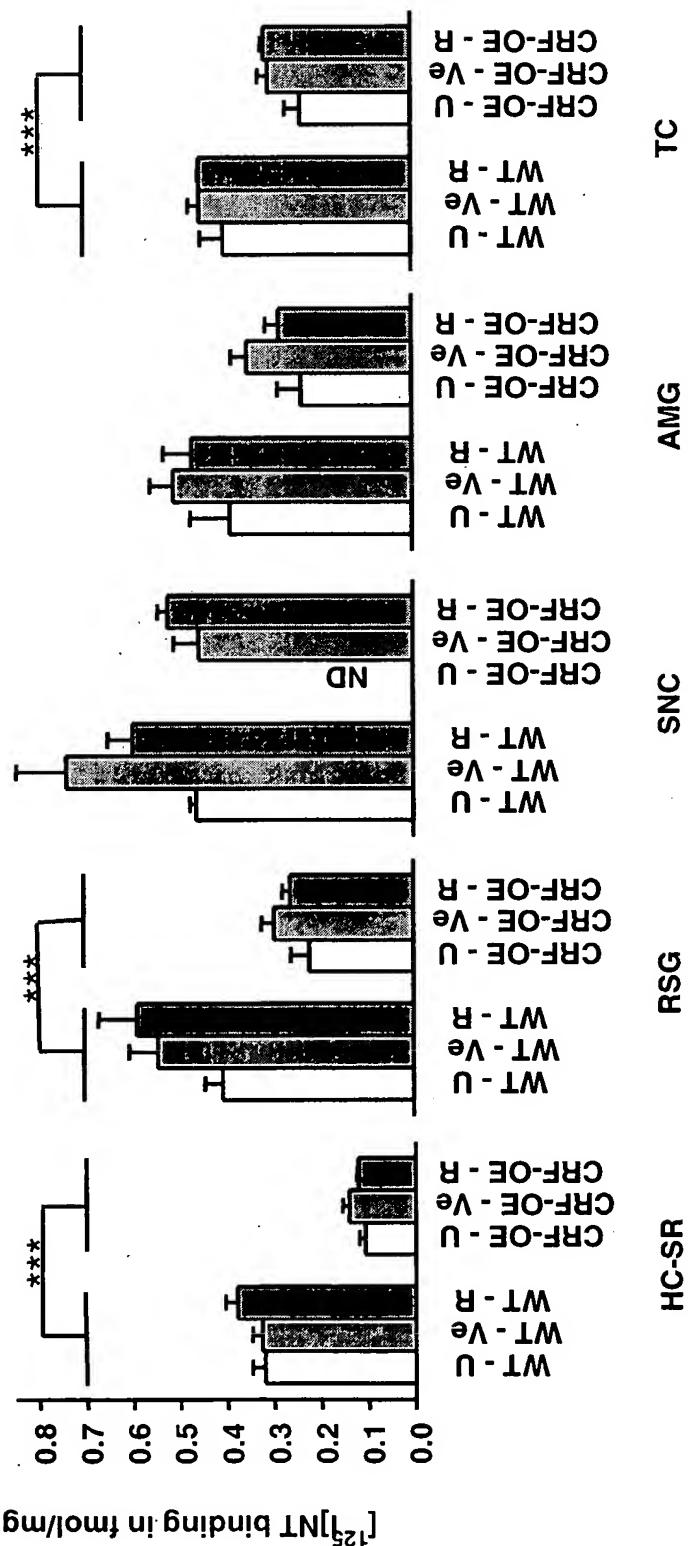


Fig.7



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